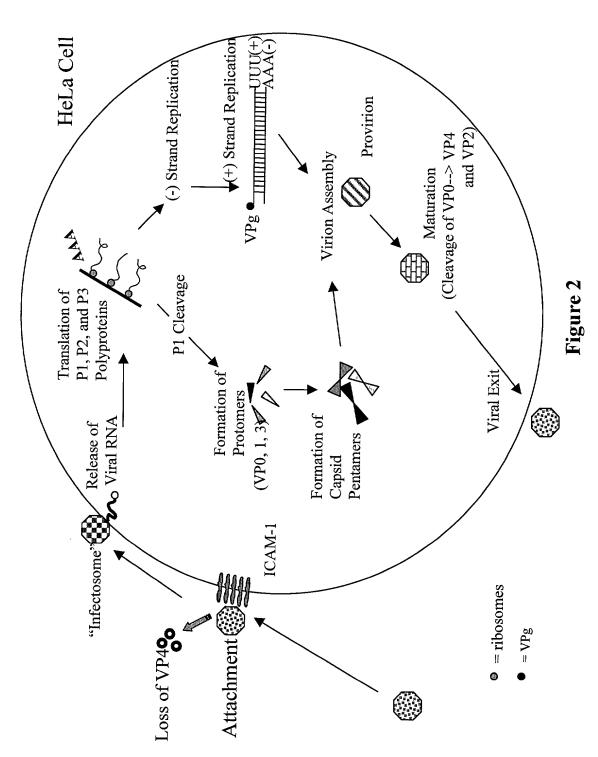
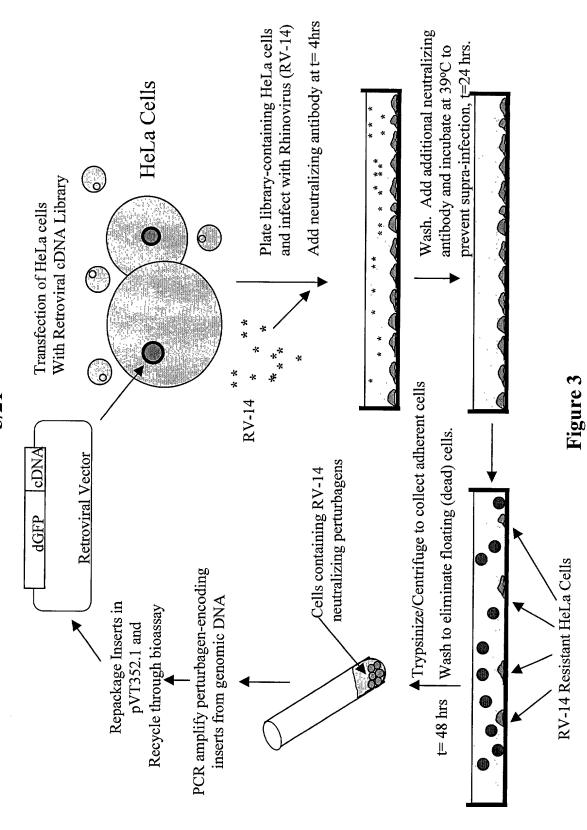


Figure 1







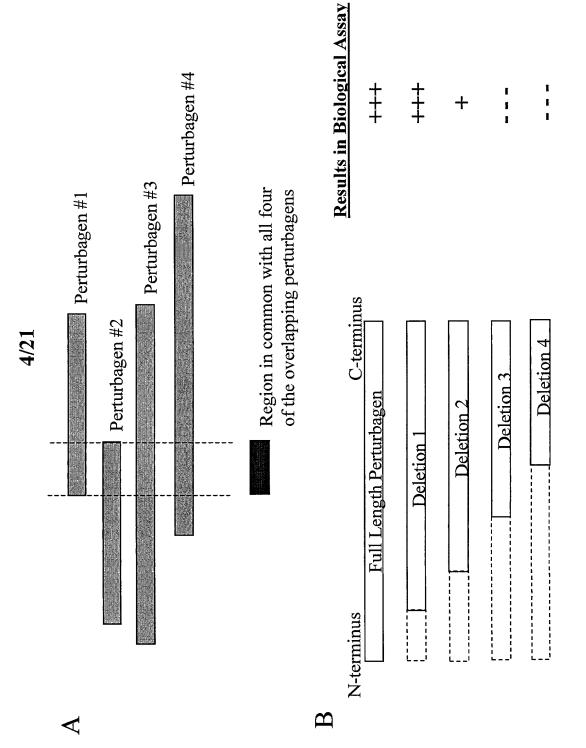


Figure 4

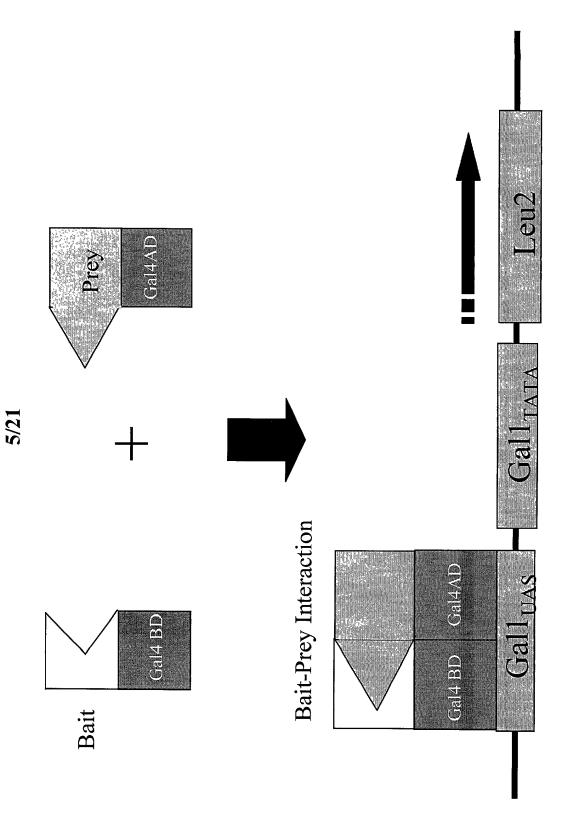


Figure 5

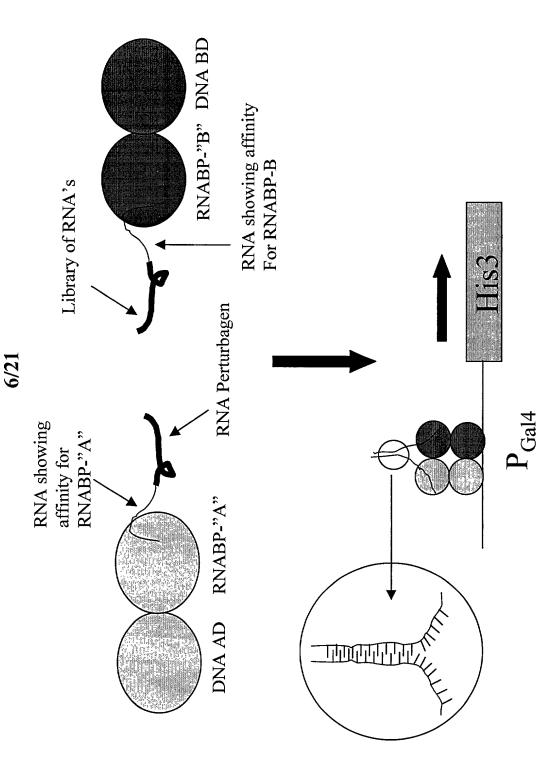
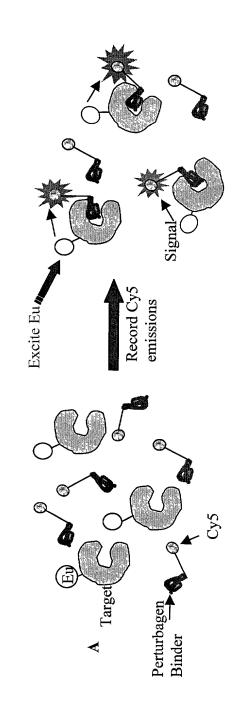
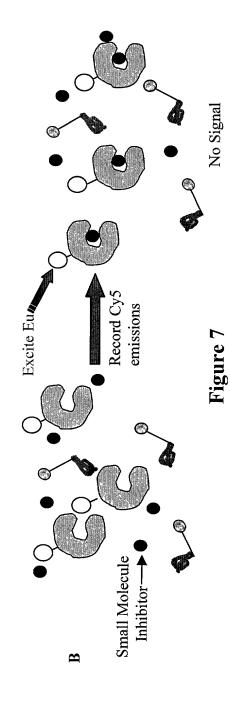


Figure 6

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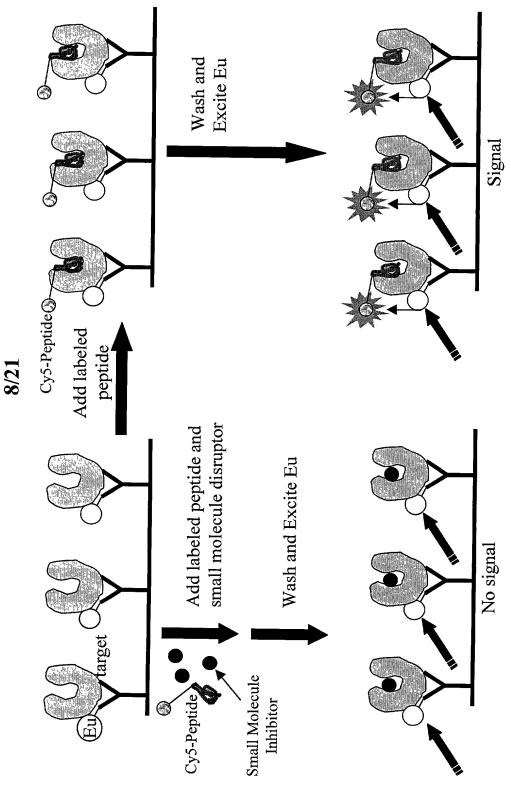


Figure 8

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Conditions for Each Cycle

Cycle 1:

24 hrs: wash + Ab K

120 hrs: harvest

24 hrs: wash + AbM

48 hrs:trypsinize/reinfect at half the moi

72 hrs: wash + Ab

96 hrs: wash + Ab

120 hrs: harvest

Cycle 2: A and B

24 hrs: wash + Ab, shift to 39° C

48 hrs: wash Ab

72 hrs: harvest

Cycle 3: A and B

24 hrs: wash + Ab, shift to 39°C 48 hrs: harvested

Note: Ab was added to all samples at 4 hrs post infection

Figure 9

Cells Surviving Each Cycle

Cycle

	1	2	3	4
352.1	0.05%	0.76%	0.07%	0.10%
Group A	0.08%	0.14%	0.17%	1.11%
Group B	<0.01%	0.22%	1.92%	6.67%

Figure 10

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DNA Sequence of cW985

5'CCGAGGCGGCCTGATAATGATGATGGTGATG ATAATGATGGTGGTGGTGAGAATAATGG ATGATGATGATGATAATGATGATGGTGG AAGATGGTGATGGTCATGATGATAGAAATGA **FGGTGGTAGGGATGATGATAATGGTGATGGT** TGGTGAGAATAATGGTGGTGATGGTGATAAT AATGATGATGGTGGATGATGGTGGTGGT **TGGTGATGATAATTATGCTACTGATGAC**
 FATGCTACTGATGACATAATGATGTGATGAT
GATGGTCGTAGTTATGCTACTGATGATA raatgatgacgctggccataatgc-3,

Amino Acid Sequence of cW985

N'- DNDDGDDNDDGGGENNGGDGDNYAT DDMMMVMIMIMMVVVRIMVVMVIIML LMT-C'

Figure 11



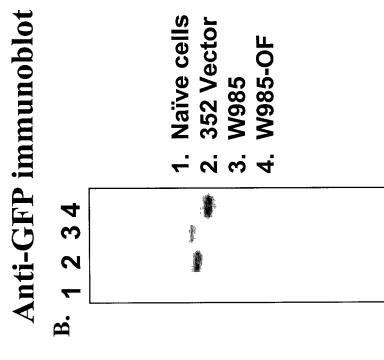


Figure 12

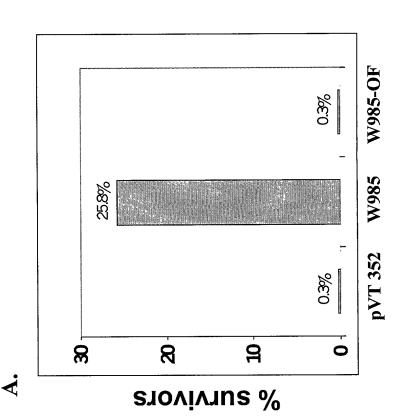


Figure 13



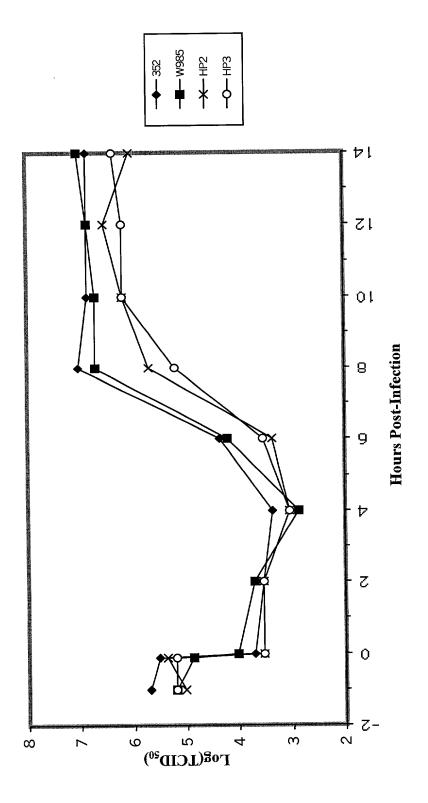


Figure 14

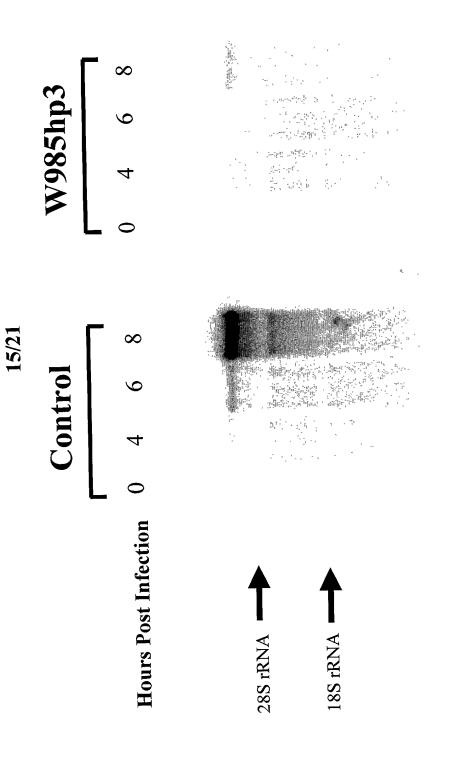


Figure 15

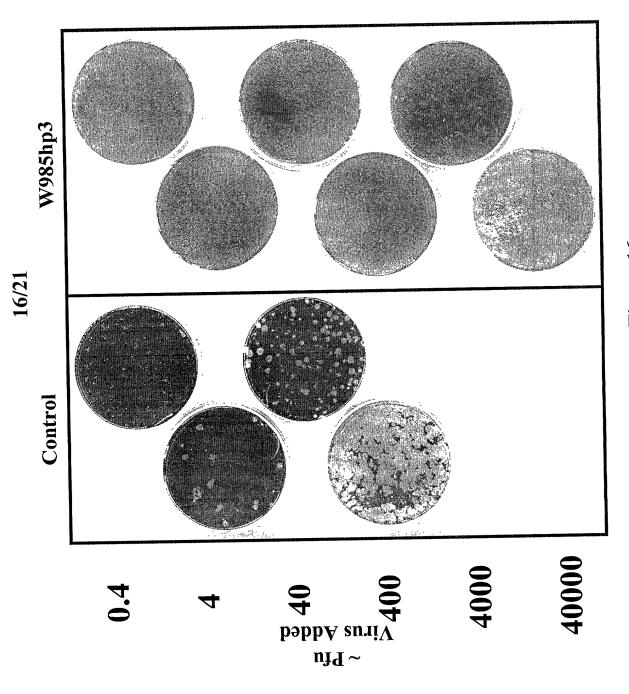
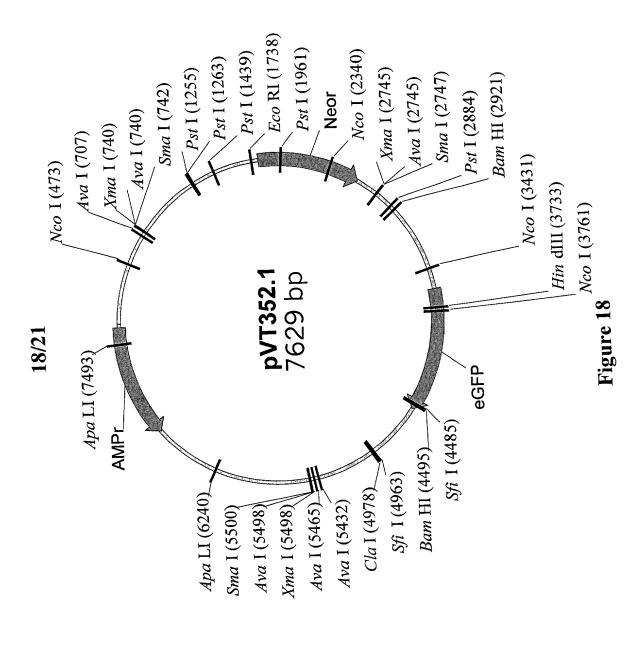


Figure 16

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3001 rh1A.F1 tcataaatggatccagAT GGGCGCTCAGGTTTCTACACA	3011 rnzB.r1 tcataaatggatccagGGCTGAGTGATTACATCACAGGTTT
3002 rh1A.R1 tcaatacagcggccgcttaATTCAATGCTGGTGCACCCTTAAGCA	3012 rh2B.R2 tcaatacagcggcttaTTGTCTTTCAATGTAAGGCACCTGA
TGA GA T	3013 rh2C.F1
3003 rh1B.F1 tcataaatggatccagTCACCCAATGTTGAGGCCTGTGGTTAT	tcataaatggatccaggchhhigh ghiggs gan a g
3004 rh1B.R1 tcaatacagcggccgcttaTTGTGGCACAA TTGACTTGGACCTT	3014 rh2C.R1 tcaatacagcggccgcttaTTGAAACAGTGTTTCTAGGGAATC
3005 rh1C.F1 tcataaatggatccagGGTTTGCCAACTA CAACTTTGCC	3015 rh3A/B.F1 tcataaatggatccagGGACCAGTGTATAAAGATTTAGAGAT
3006 rh1C.R1 tcaatacagcggccgcttaTTCAGTGAGTGCAACAGTCTGTG	3016 rh3A/B.R1 tcaatacagcggccgcttaTTGCACAACAACTGGGCGTAAAGT
3007 rh1D.F1 tcataaatggatccagGGCTTA GGTGATGAATTA GAAGAA GT	3017 rh3C.F1 tcataaatggatccagGGACCAAACACAGAATTTGCA CTAT C
3008 rh1D.R2 tcaatacagcggccgcttaATAGGATTTAATGTCACCTTTCCTCT 3009 rh2A.F1	3018 rh3C.R2 tcaatacagcggccgcttaTTGTTTCTCTACAAATAT TGTTTTTAAGTTGAGC
3010 rh2A.Rl	3019 rh3D.F1 acataaatgcggccgcaaGGCCAAGTAATAGCTAGACATAAG
	3020 rh3D.R1 tcaatacagcggccgcttaAAAGAGGTCCAACCAGCGCCTCCTAA

Figure 17



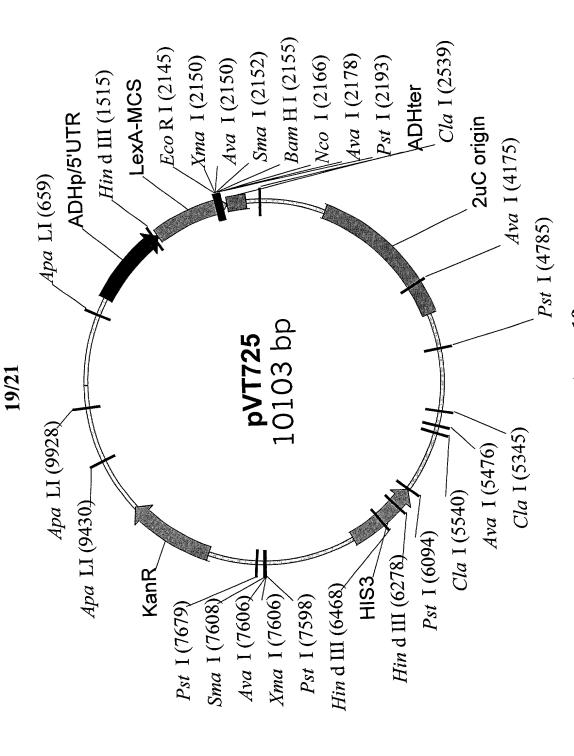


Figure 19

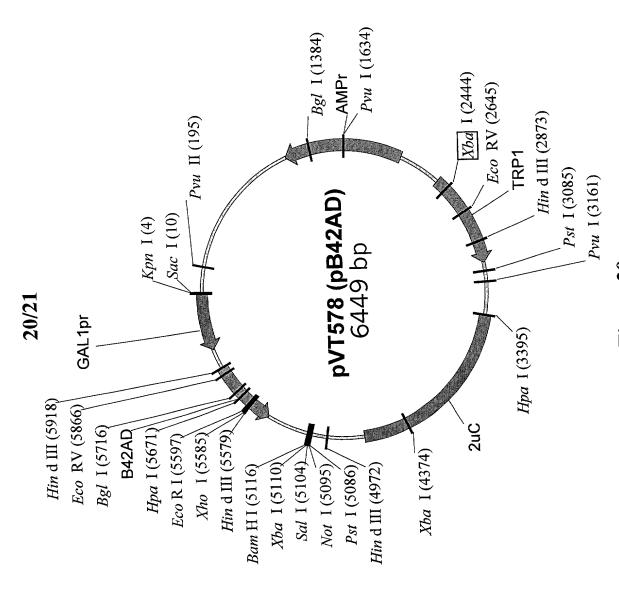


Figure 20

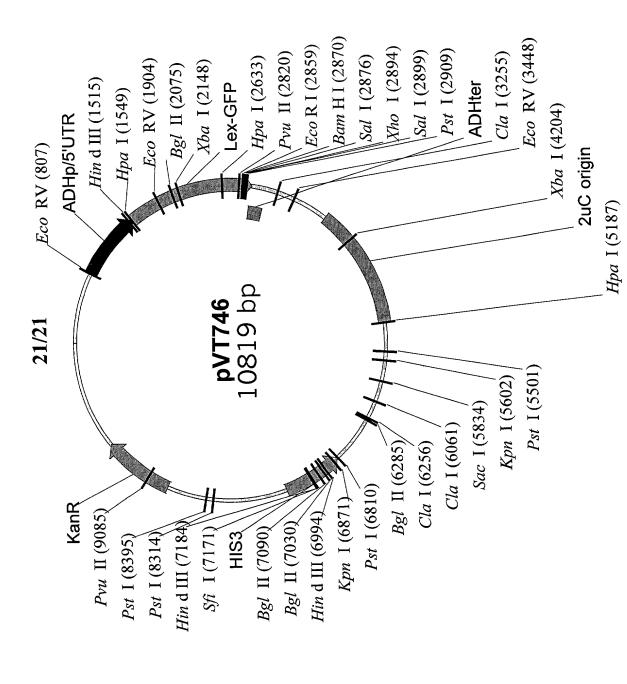


Figure 2]